SEQUENCE LISTING

```
<110> QIN, MINMIN
            CHIN, WIA-PAN
            CHEN, LIN
            FITZPATRICK, PAUL A.
            HENSTRAND, JOHN M.
            WENDTH, DAN J.
            ZECHERLE, GARY N.
            STARR, CHRISTOPEHER M.
            KAKKIS, EMIL D.
      <120> METHODS FOR PRODUCING AND PURIFYING RECOMBINANT ALPHA-L-
            IDURONIDASE
      <130> 008000051CNUS02
      <140> TO BE ASSIGNED
      <141> 2001-11-13
      <150> 09/711,202
      <151> 2000-11-09
      <150> 09/439,923
      <151> 1999-11-12
      <160> 2
      <170> FastSEQ for Windows Version 3.0
      <210> 1
      <211> 6200
      <212> DNA
      <213> Homo sapiens
      <220>
      <221> CDS
      <222> (1558)...(3510)
      <400> 1
gacggategg gagatetece gatecectat ggtegactet cagtacaate tgetetgatg
ccgcatagtt aagccagtat ctgctccctg cttgtgtgtt ggaggtcgct gagtagtgcg
                                                                       120
cqaqcaaaat ttaagctaca acaaggcaag gcttgaccga caattgcatg aagaatctgc
                                                                       180
ttagggttag gcgttttgcg ctgcttcgcg atgtacgggc cagatatacg cgttgacatt
                                                                       240
gattattgac tagttattaa tagtaatcaa ttacggggtc attagttcat agcccatata
                                                                       300
tggagttccg cgttacataa cttacggtaa atggcccgcc tggctgaccg cccaacgacc
                                                                       360
                                                                       420
cccgcccatt gacgtcaata atgacgtatg ttcccatagt aacgccaata gggactttcc
                                                                       480
attgacgtca atgggtggac tatttacggt aaactgccca cttggcagta catcaagtgt
                                                                       540
atcatatgcc aagtacgccc cctattgacg tcaatgacgg taaatggccc gcctggcatt
atgcccagta catgacctta tgggactttc ctacttggca gtacatctac gtattagtca
                                                                       600
tcgctattac catggtgatg cggttttggc agtacatcaa tgggcgtgga tagcggtttg
                                                                       660
actcacgggg atttccaagt ctccacccca ttgacgtcaa tgggagtttg ttttggcacc
                                                                       720
aaaatcaacg ggactttcca aaatgtcgta acaactccgc cccattgacg caaatgggcg
                                                                       780
gtaggcgtgt acggtgggag gtctatataa gcagagctct ctggctaact agagaaccca
                                                                       840
ctgcttaact ggcttatcga aattaatacg actcactata gggagaccca agcttcgcag
                                                                       900
aatteetgeg getgetacag tgtgteeage gteetgeetg getgtgetga gegetggaac
                                                                       960
agtggcgcat cattcaagtg cacagttacc catcctgagt ctggcacctt aactggcaca
                                                                      1020
attgccaaag tcacaggtga gctcagatgc ataccaggac attgtatgac gttccctgct
                                                                      1080
cacatgootg otttottoot ataatacaga tgotcaacta actgotcatg toottatato
                                                                      1140
acagagggaa attggagcta tctgaggaac tgcccagaag ggaagggcag aggggtcttg
                                                                      1200
```

character at additional activities at activities at activities at a construction of the construction of th												1260 1320 1380 1440 1500 1560				
cgt Arg	ccc Pro	ctg Leu	cgc Arg 5	ccc	cgc Arg	gcc Ala	gcg Ala	ctg Leu 10	ctg Leu	gcg Ala	ctc Leu	ctg Leu	gcc Ala 15	tcg Ser	ctc Leu	1608
ctg Leu	gcc Ala	gcg Ala 20	ccc	ccg Pro	gtg Val	gcc Ala	ccg Pro 25	gcc Ala	gag Glu	gcc Ala	ccg Pro	cac His 30	ctg Leu	gtg Val	cat His	1656
gtg Val	gac Asp 35	gcg Ala	gcc Ala	cgc Arg	gcg Ala	ctg Leu 40	tgg Trp	ccc Pro	ctg Leu	cgg Arg	cgc Arg 45	ttc Phe	tgg Trp	agg Arg	agc Ser	1704
aca Thr 50	ggc Gly	ttc Phe	tgc Cys	ccc Pro	ccg Pro 55	ctg Leu	cca Pro	cac His	agc Ser	cag Gln 60	Ala	gac Asp	cag Gln	tac Tyr	gtg Val 65	1752
ctc Leu	agc Ser	tgg Trp	gac Asp	cag Gln 70	cag Gln	ctc Leu	aac Asn	ctc Leu	gcc Ala 75	tat Tyr	gtg Val	ggc Gly	gcc Ala	gtc Val 80	PIO	1800
cac His	cgc Arg	ggc Gly	atc Ile 85	aag Lys	cag Gln	gtc Val	cgg Arg	acc Thr 90	His	tgg Trp	ctg Leu	ctg Leu	gag Glu 95	ctt Leu	gtc Val	1848
acc Thr	acc Thr	agg Arg 100	ggg Gly	tcc Ser	act Thr	gga Gly	cgg Arg 105	ggc Gly	ctg Leu	agc Ser	tac Tyr	aac Asn 110	ttc Phe	acc Thr	cac His	1896
ctg Leu	gac Asp 115	Gly ggg	tac Tyr	ctg Leu	gac Asp	ctt Leu 120	ctc Leu	agg Arg	gag Glu	aac Asn	cag Gln 125	ctc Leu	Gly 999	ttt Phe	gag Glu	1944
ctg Leu 130	atg Met	ggc Gly	agc Ser	gcc Ala	tcg Ser 135	ggc Gly	cac His	ttc Phe	act Thr	gac Asp 140	ttt Phe	gag Glu	gac Asp	aag Lys	Gln 145	1992
cag Gln	gtg Val	ttt Phe	gag Glu	tgg Trp 150	aag Lys	gac Asp	ttg Leu	gtc Val	tcc Ser 155	agc Ser	ctg Leu	gcc Ala	agg Arg	aga Arg 160	tac Tyr	2040
atc Ile	ggt Gly	agg Arg	tac Tyr 165	Gly	ctg Leu	gcg Ala	cat His	gtt Val 170	tcc Ser	aag Lys	tgg Trp	aac Asn	Phe	GIU	acg Thr	2088
tgg Trp	aat Asn	gag Glu 180	cca Pro	gac Asp	cac His	cac His	gac Asp 185	ttt Phe	gac Asp	aac Asn	gtc Val	ser 190	Met	aco Thr	atg Met	2136
caa Gln	ggc Gly 195	ttc Phe	ctg Leu	aac Asn	tac Tyr	tac Tyr 200	gat Asp	gcc Ala	tgc Cys	tcg Ser	gag Glu 205	GIY	ctg Lev	cgo Arg	gcc g Ala	2184

					-			٠.									
7	gcc Ala 210	agc Ser	ccc Pro	gcc Ala	ctg Leu	cgg Arg 215	ctg Leu	gga Gly	ggc Gly	ccc Pro	ggc Gly 220	gac Asp	tcc Ser	ttc Phe	cac His	acc Thr 225	2232
]	ca Pro	ccg Pro	cga Arg	tcc Ser	ccg Pro 230	ctg Leu	agc Ser	tgg Trp	ggc Gly	ctc Leu 235	ctg Leu	cgc Arg	cac His	Cye Cye	cac His 240	gac Asp	2280
9.	ggt 31y	acc Thr	aac Asn	ttc Phe 245	ttc Phe	act Thr	999 Gly	gag Glu	gcg Ala 250	ggc Gly	gtg Val	cgg Arg	ctg Leu	gac Asp 255	tac Tyr	atc Ile	2328
	tċc Ser	ctc Leu	cac His 260	agg Arg	aag Lys	ggt Gly	gcg Ala	cgc Arg 265	agc Ser	tcc Ser	atc Ile	tcc Ser	atc Ile 270	ctg Leu	gag Glu	cag Gln	2376
	gag Glu	aag Lys 275	gtc Val	gtc Val	gcg Ala	cag Gln	cag Gln 280	atc Ile	cgg	cag Gln	ctc Leu	ttc Phe 285	ccc Pro	aag Lys	ttc Phe	gcg Ala	2424
	gac Asp 290	acc Thr	ccc Pro	att Ile	tac Tyr	aac Asn 295	gac Asp	gag Glu	gcg Ala	gac Asp	ccg Pro 300	ctg Leu	gtg Val	ggc Gly	tgg Trp	ser 305	2472
	ctg Leu	cca Pro	cag Gln	ccg Pro	tgg Trp 310	Arg	gcg Ala	gac Asp	gtg Val	acc Thr 315	Tyr	gcg Ala	gcc Ala	atg Met	gtg .Va] 320	g gtg Val	2520
	aag Lys	gtc Val	atc Ile	gcg Ala 325	cag Gln	cat His	cag Gln	aac Asn	ctg Leu 330	Leu	ctg Leu	gcc Ala	aac Asn	acc Thr 335	. 1111	c tcc Ser	2568
	gcc Ala	ttc Phe	ccc Pro	Tyr	gcg Ala	ctc Leu	ctg Leu	agc Ser 345	Asn	gac Asp	aat Asn	gcc Ala	ttc Phe 350	Let	g ago 1 Sei	tac r Tyr	2616
	cac His	ccg Pro 355	His	ccc Pro	ttc Phe	gcg Ala	cag Gln 360	Arg	acg Thr	ctc Leu	acc	gcg Ala 365	a Arg	tto Phe	c cae	g gtc n _. Val	2664
	aac Asn 370	Asn	acc Thr	cgc Arg	ccg Pro	ccg Pro 375	His	gtg Val	cag Gln	ctg Leu	ı Lev	g cgc L Arg	д ГА:	g cc	g gt o Va	g ctc l Leu 385	2712
•	acg Thr	gcc Ala	atg Met	: Gly	ctg Leu 390	Leu	gcg	ctg Leu	ctg Lev	gat Asp 395	Glu	g gag 1 Glv	g cag u Glr	g ct	c tg u Tr 40	g gcc p Ala 0	2760
	gaa Glu	gtg Val	tcg Sei	cag Gln 405	Ala	ggg Gly	acc Thr	gtc Val	cto Lev	ı Asp	ago Sei	c aac	c cad	c ac s Th 41	T va	g ggc 1 Gly	2808
	gto Val	cto Lei	g gco 1 Ala 420	a Ser	gcc Ala	cac His	cgc Arg	200 Pro 425	GT1	g ggo n Gly	e ccg	g gc	c gad a Asj 43	р мт	c tg a Tr	g cgc	2856
	gcc	gcg Ala 435	a Val	g ctg Leu	ato Ile	tac Tyr	gcg Ala	Ser	gad Asp	gad Asp	c acc	c cg r Ar	g AT	c ca a Hi	c co s Pr	c aac	2904

cgc Arg 450	agc Ser	gtc Val	gcg Ala	gtg Val	acc Thr 455	ctg Leu	cgg Arg	ctg Leu	cgc Arg	999 Gly 460	gtg Val	ccc Pro	ccc Pro	ggc	ccg Pro 465	2952
ggc Gly	ctg Leu	gtc Val	tac Tyr	gtc Val 470	acg Thr	cgc Arg	tac Tyr	ctg Leu	gac Asp 475	aac Asn	ggg ggg	ctc Leu	tgc Cys	agc Ser 480	ccc Pro	.3000
gac Asp	ggc Gly	gag Glu	tgg Trp 485	cgg Arg	cgc Arg	ctg Leu	ggc Gly	cgg Arg 490	ccc Pro	gtc Val	ttc Phe	ccc Pro	acg Thr 495	gca Ala	gag Glu	3048
Gln	ttc Phe	Arg 500	Arg	*	Arg	Ala	Ala	Glu 505	Asp	Pro	Val	Ala	Ala 510	Ala	Pro	3096
Arg	ccc Pro	Leu 515	Pro	Ala	Gly	Gly	Arg 520	Leu	Arg	Leu	Arg	Pro 525	Ala	Leu	Arg	3144
Leu	530	Ser	Leu	Leu	Leu	Val 535	His	Val	Cys	Ala	Arg 540	Pro	Glu	Lys	Pro.	3192
Pro 545		Gln	Val	Thr	Arg 550	Leu ·	Arg	Ala	Leu	Pro .555	Leu	Thr	Gln	Gly	Gln 560	3240
Leu	Val	Leu	Val	Trp 565	Ser	Asp	Glu	His	Val 570	Gly	Ser	Lys	Cys	.Leu 575		3288
Thr	Tyr	Glu	Ile 580	Gln	Phe	Ser	Gln	Asp 585	Gly	Lys	Ala	Tyr	Thr 590	Pro	gtc Val	3336
Ser	Arg	Lys 595	Pro	Ser	Thr	Phe	Asn 600	Leu	Phe	Val	Phe	Ser 605	Pro	Asp	aca Thr	3384
Gly	Ala 610	Val	Ser	Gly	Ser	Tyr 615	Arg	Val	Arg	Ala	Leu 620	Asp	Туз	Trp	gcc Ala	3432
Arg 625	Pro	Gly	Pro	Phe	Ser 630	Asp	Pro	Val	Pro	Tyr 635	Leu	Glu	. Va]	l Pro	gtg Val 640	3480
Pro	aga Arg	Gly	Pro	Pro 645	Ser	Pro	Gly	Asn	Pro 650			•				3530
ccc aaa gcc gcc agg tag	atca aaaa cgggg ttcta tgcca gtgte caata	aaa gga agt act cat	cctti aaaaa tccad tgcca cccad tctai	tgcaa aaaaa ctagt agcca ctgto ttcto	at a	tatti aaaa tagag tgttg tcci gggtg	tta aaaa gggc gttt gtaat gggg gegg	t at aa aa c cg cc a aa t gg	ttta aaaa ttta cctc atga ggca gctc	aaaa aaacc cccc ggaa ggac tatg	aaa aaa cgc gtg att agc	aaaa tgat cctt gcat aagg	aaa cag cct cct cgc 1999	aaaa aatt cctc tgac attc agga cgga	etgeed aaaaa ceetge eectge gtetga attgge aaagaa aaattg	3650 3710 3770 3830 3830 3890 3890 3950 300
								-	46-							
		•		,										•		ā

```
tatccgctca caattccaca caacatacga gccggaagca taaagtgtaa agcctggggt
                                                                    4130
gectaatgag tgagetaact cacattaatt gegttgeget cactgeeege tttecagteg
                                                                    4190
ggaaacctgt cgtgccagct gcattaatga atcggccaac gcgcggggag aggcggtttg
                                                                    4250
cgtattgggc gctcttccgc ttcctcgctc actgactcgc tgcgctcggt cgttcggctg
                                                                    4310
cggcgagcgg tatcagctca ctcaaaggcg gtaatacggt tatccacaga atcaggggat
                                                                    4370
aacgcaggaa agaacatgtg agcaaaaggc cagcaaaagg ccaggaaccg taaaaaggcc
                                                                    4430
gegttgetgg egttttteca taggeteege eeeeetgaeg ageateacaa aaategaege
                                                                    4490
tcaagtcaga ggtggcgaaa cccgacagga ctataaagat accaggcgtt tccccctgga
                                                                     4550
ageteceteg tgegetetee tgtteegace etgeegetta eeggataeet gteegeettt
                                                                     4610
ctcccttcgg gaagcgtggc gctttctcaa tgctcacgct gtaggtatct cagttcggtg
                                                                     4670
                                                                     4730
taggtcgttc gctccaagct gggctgtgtg cacgaacccc ccgttcagcc cgaccgctgc
gccttatccg gtaactatcg tcttgagtcc aacccggtaa gacacgactt atcgccactg
                                                                     4790
gcagcagcca ctggtaacag gattagcaga gcgaggtatg taggcggtgc tacagagttc
                                                                     4850
ttgaagtggt ggcctaacta cggctacact agaaggacag tatttggtat ctgcgctctg
                                                                     4910
ctgaagccag ttaccttcgg aaaaagagtt ggtagctctt gatccggcaa acaaaccacc
                                                                     4970
gctggtagcg gtggtttttt tgtttgcaag cagcagatta cgcgcagaaa aaaaggatct
                                                                     5030
caagaagatc ctttgatctt ttctacgggg tctgacgctc agtggaacga aaactcacgt
                                                                     5090
taagggattt tggtcatgag attatcaaaa aggatcttca cctagatcct tttaaattaa
                                                                     5150
aaatgaagtt ttaaatcaat ctaaagtata tatgagtaaa cttggtctga cagttaccaa
                                                                     5210
tgcttaatca gtgaggcacc tatctcagcg atctgtctat ttcgttcatc catagttgcc
                                                                     5270
tgactccccg tcgtgtagat aactacgata cgggagggct taccatctgg ccccagtgct
                                                                     5330
gcaatgatac cgcgagaccc acgctcaccg gctccagatt tatcagcaat aaaccagcca
                                                                     5390
gccggaaggg ccgagcgcag aagtggtcct gcaactttat ccgcctccat ccagtctatt
                                                                     5450
aattgttgcc gggaagctag agtaagtagt tcgccagtta atagtttgcg caacgttgtt
                                                                     5510
gccattgcta caggcatcgt ggtgtcacgc tcgtcgtttg gtatggcttc attcagctcc
                                                                     5570 ·
ggttcccaac gatcaaggcg agttacatga tcccccatgt tgtgcaaaaa agcggttagc
                                                                     5630
teetteggte eteegategt tgteagaagt aagttggeeg cagtgttate acteatggtt
                                                                     5690
atggcagcac tgcataattc tcttactgtc atgccatccg taagatgctt ttctgtgact
                                                                     5750
ggtgagtact caaccaagtc attctgagaa tagtgtatgc ggcgaccgag ttgctcttgc
                                                                     5810
ceggegteaa taegggataa taeegegeea catageagaa etttaaaagt geteateatt
                                                                     5870
ggaaaacgtt cttcggggcg aaaactctca aggatcttac cgctgttgag atccagttcg
                                                                     5930
atgtaaccca ctcgtgcacc caactgatct tcagcatctt ttactttcac cagcgtttct
                                                                     5990
                                                                     6050
gggtgagcaa aaacaggaag gcaaaatgcc gcaaaaaagg gaataagggc gacacggaaa
tgttgaatac tcatactctt cctttttcaa tattattgaa gcatttatca gggttattgt
                                                                     6110
ctcatgagcg gatacatatt tgaatgtatt tagaaaaata aacaaatagg ggttccgcgc
                                                                     6170
                                                                      6200
acatttcccc gaaaagtgcc acctgacgtc
```

<210> 2 <211> 650 <212> PRT <213> Homo sapiens

Met Arg Pro Leu Arg Pro Arg Ala Ala Leu Leu Ala Leu Leu Ala Ser 10 Leu Leu Ala Ala Pro Pro Val Ala Pro Ala Glu Ala Pro His Leu Val 25 20 His Val Asp Ala Ala Arg Ala Leu Trp Pro Leu Arg Arg Phe Trp Arg 40 Ser Thr Gly Phe Cys Pro Pro Leu Pro His Ser Gln Ala Asp Gln Tyr 55 Val Leu Ser Trp Asp Gln Gln Leu Asn Leu Ala Tyr Val Gly Ala Val 70 Pro His Arg Gly Ile Lys Gln Val Arg Thr His Trp Leu Leu Glu Leu 90 85 Val Thr Thr Arg Gly Ser Thr Gly Arg Gly Leu Ser Tyr Asn Phe Thr 110 105 100 His Leu Asp Gly Tyr Leu Asp Leu Leu Arg Glu Asn Gln Leu Gly Phe 120 125 Glu Leu Met Gly Ser Ala Ser Gly His Phe Thr Asp Phe Glu Asp Lys 140 135

```
Gln Gln Val Phe Glu Trp Lys Asp Leu Val Ser Ser Leu Ala Arg Arg
                                        155
Tyr Ile Gly Arg Tyr Gly Leu Ala His Val Ser Lys Trp Asn Phe Glu
                                    170
                165
Thr Trp Asn Glu Pro Asp His His Asp Phe Asp Asn Val Ser Met Thr
                                185
Met Gln Gly Phe Leu Asn Tyr Tyr Asp Ala Cys Ser Glu Gly Leu Arg
                                                205
                            200
        195
Ala Ala Ser Pro Ala Leu Arg Leu Gly Gly Pro Gly Asp Ser Phe His
                                            220
                        215
Thr Pro Pro Arg Ser Pro Leu Ser Trp Gly Leu Leu Arg His Cys His
                                        235
                    230
Asp Gly Thr Asn Phe Phe Thr Gly Glu Ala Gly Val Arg Leu Asp Tyr
                                    250
                245
Ile Ser Leu His Arg Lys Gly Ala Arg Ser Ser Ile Ser Ile Leu Glu
                                265
            260
Gln Glu Lys Val Val Ala Gln Gln Ile Arg Gln Leu Phe Pro Lys Phe
                            280
                                                285
        275
Ala Asp Thr Pro Ile Tyr Asn Asp Glu Ala Asp Pro Leu Val Gly Trp
                        295
                                            300
    290
Ser Leu Pro Gln Pro Trp Arg Ala Asp Val Thr Tyr Ala Ala Met Val
                                        315
                    310
Val Lys Val Ile Ala Gln His Gln Asn Leu Leu Leu Ala Asn Thr Thr
                                    330
                325
Ser Ala Phe Pro Tyr Ala Leu Leu Ser Asn Asp Asn Ala Phe Leu Ser
                                                    350
                                345
            340
Tyr His Pro His Pro Phe Ala Gln Arg Thr Leu Thr Ala Arg Phe Gln
                                                365
                            360
Val Asn Asn Thr Arg Pro Pro His Val Gln Leu Leu Arg Lys Pro Val
                                            380
                        375
Leu Thr Ala Met Gly Leu Leu Ala Leu Leu Asp Glu Glu Gln Leu Trp
                                        395
                    390
Ala Glu Val Ser Gln Ala Gly Thr Val Leu Asp Ser Asn His Thr Val
                                                       415
                                    410
                405
Gly Val Leu Ala Ser Ala His Arg Pro Gln Gly Pro Ala Asp Ala Trp
                                                    430
                                425
            420
Arg Ala Ala Val Leu Ile Tyr Ala Ser Asp Asp Thr Arg Ala His Pro
                                                445
                            440
Asn Arg Ser Val Ala Val Thr Leu Arg Leu Arg Gly Val Pro Pro Gly
                                             460
                        455
Pro Gly Leu Val Tyr Val Thr Arg Tyr Leu Asp Asn Gly Leu Cys Ser
                                       . 475
                    470
Pro Asp Gly Glu Trp Arg Arg Leu Gly Arg Pro Val Phe Pro Thr Ala
                                     490
                485
Glu Gln Phe Arg Arg Ala Ala Glu Asp Pro Val Ala Ala Pro
                                505
Arg Pro Leu Pro Ala Gly Gly Arg Leu Arg Leu Arg Pro Ala Leu Arg
                            520
Leu Pro Ser Leu Leu Leu Val His Val Cys Ala Arg Pro Glu Lys Pro
                        535
Pro Gly Gln Val Thr Arg Leu Arg Ala Leu Pro Leu Thr Gln Gly Gln
                                        555
                    550
Leu Val Leu Val Trp Ser Asp Glu His Val Gly Ser Lys Cys Leu Trp
                                    570
                                                         575
                565
Thr Tyr Glu Ile Gln Phe Ser Gln Asp Gly Lys Ala Tyr Thr Pro Val
                                                     590
                                585
Ser Arg Lys Pro Ser Thr Phe Asn Leu Phe Val Phe Ser Pro Asp Thr
                                                 605
                            600
        595
Gly Ala Val Ser Gly Ser Tyr Arg Val Arg Ala Leu Asp Tyr Trp Ala
                                             620
                        615
Arg Pro Gly Pro Phe Ser Asp Pro Val Pro Tyr Leu Glu Val Pro Val
```

625 630 635 640

Pro Arg Gly Pro Pro Ser Pro Gly Asn Pro
645 650